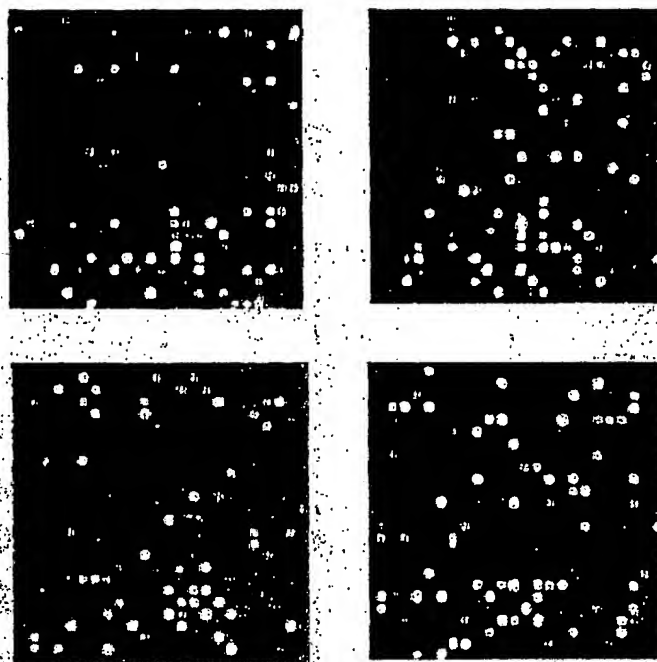


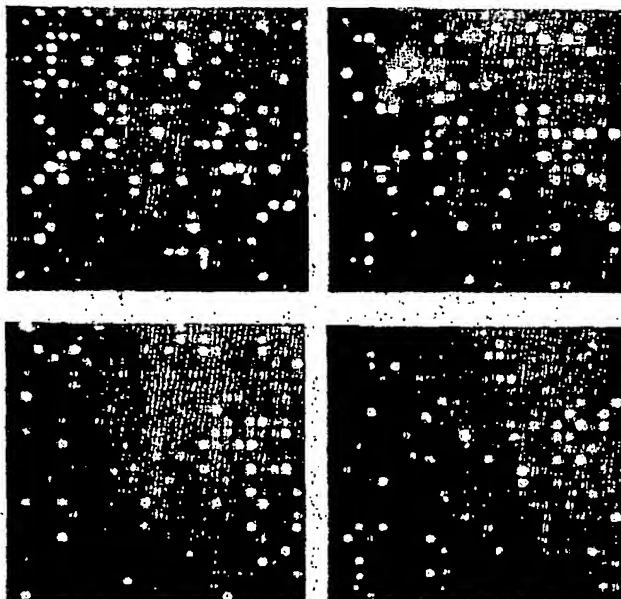
**Expression Profiling of Microdissected Tumors
Using Genentech Microarrays**



Probe generated from ~1-5 ng of total RNA (~10 - 50 pg mRNA / polyA+RNA) from a microdissected colon tumor, raw data, using amplification and probe labelling protocol.

FIG. 1

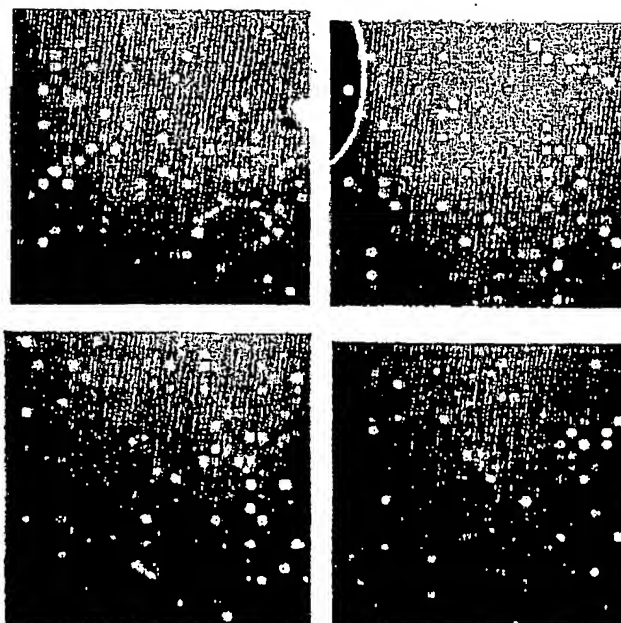
Expression Profiling of RNA from
Paraffin-Embedded Tissue Using
Genentech Microarrays



Probe generated from 5 ug total RNA,
adult liver, fresh frozen sample,
Genentech probe protocol

FIG.-2B

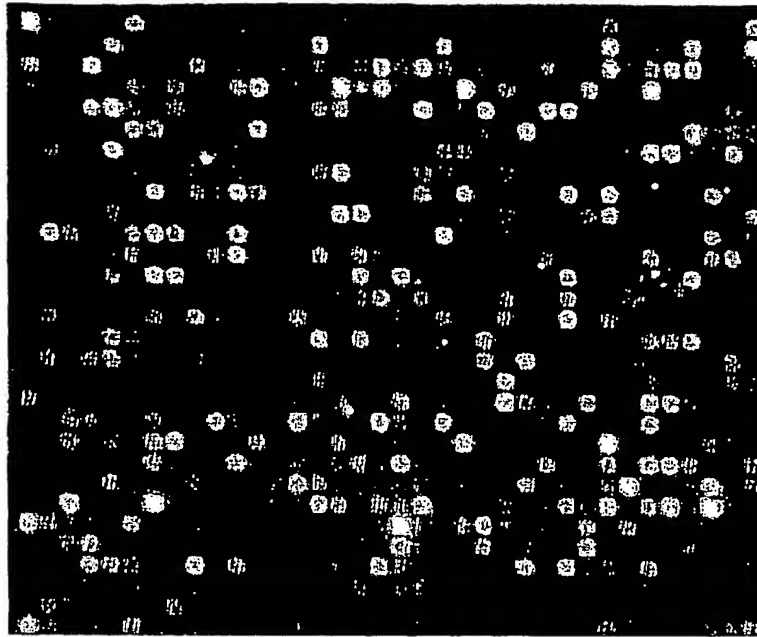
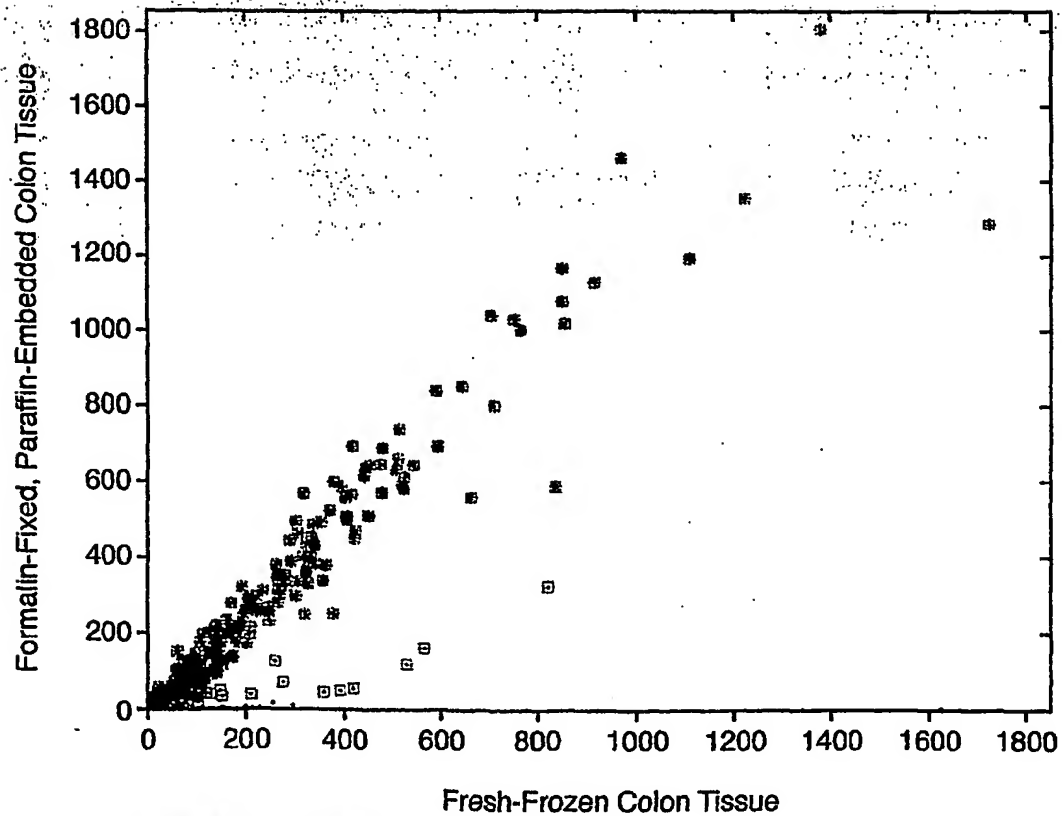
Expression Profiling of RNA from
Paraffin-Embedded Tissue Using
Genentech Microarrays



Probe generated from 5 ug total RNA, isolated
from formalin-fixed paraffin-embedded liver
tissue, Genentech probe protocol

FIG.-2A

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**FIG. 2C****FIG. 2D**

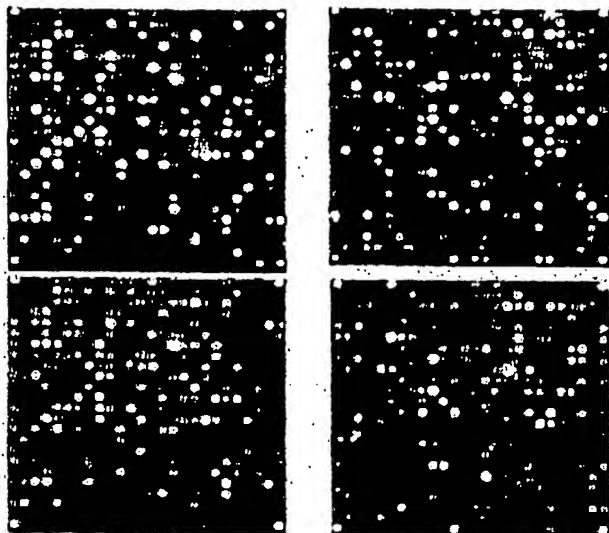
SUBSTITUTE SHEET (RULE 26)

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4/5

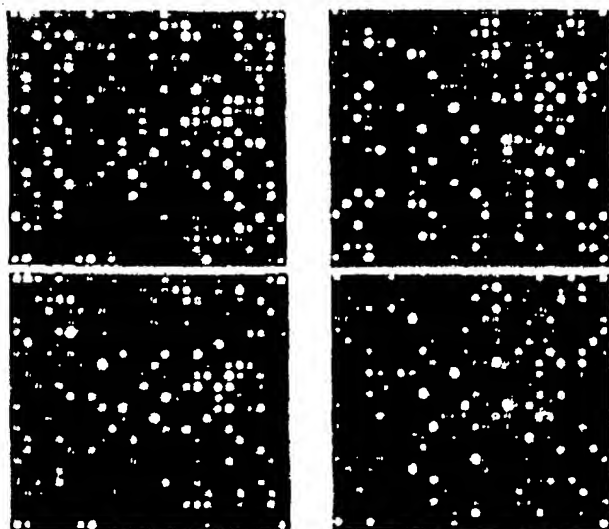
Analysis of Gene Expression Using Genetech Microarrays



Example 2496 gene microarray hybridized with a probe generated from 2.5 ug total RNA (~10 - 30 ng mRNA / polyA+RNA) from a breast tumor vs. 2.5 ug total RNA from an epithelial tissue RNA pool reference sample. The raw data from both fluorochromes for 4 subarrays are shown (Alexa-546 labelled epithelial pool probes).

FIG. 3B

Analysis of Gene Expression Using Genetech Microarrays



Example 2496 gene microarray hybridized with a probe generated from 2.5 ug total RNA (~10 - 30 ng mRNA / polyA+RNA) from a breast tumor vs. 2.5 ug total RNA from an epithelial tissue RNA pool reference sample. The raw data from both fluorochromes for 4 subarrays are shown (Alexa-488 labelled tumor probes).

FIG. 3A

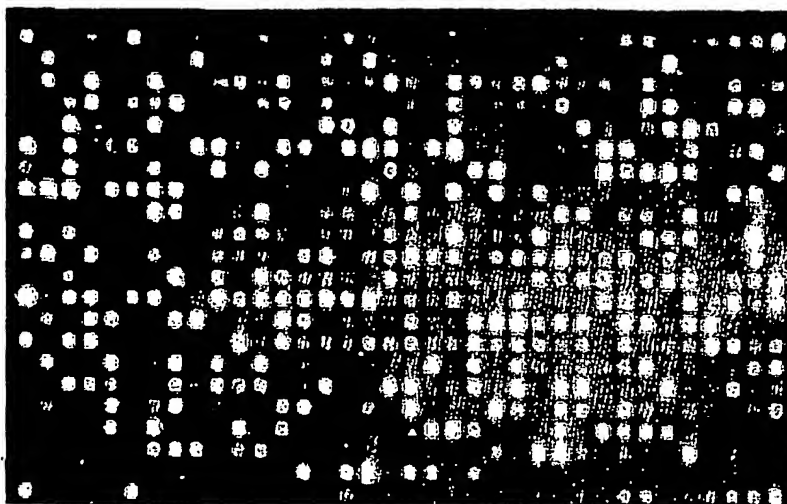


FIG. 4A

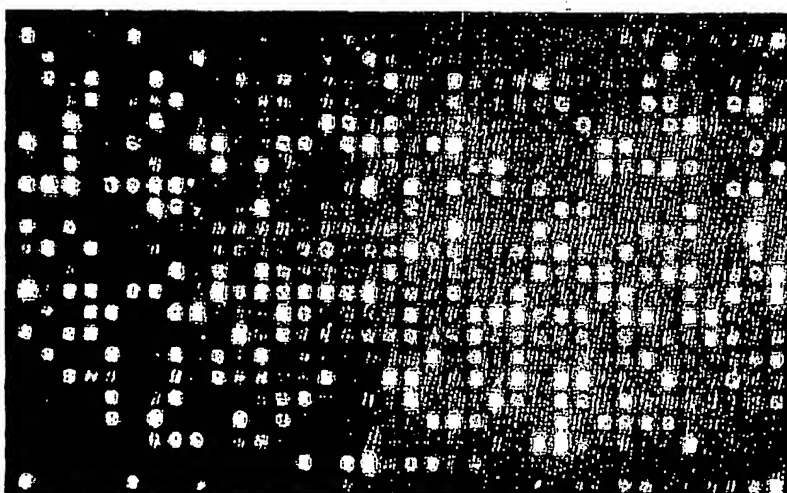


FIG. 4B

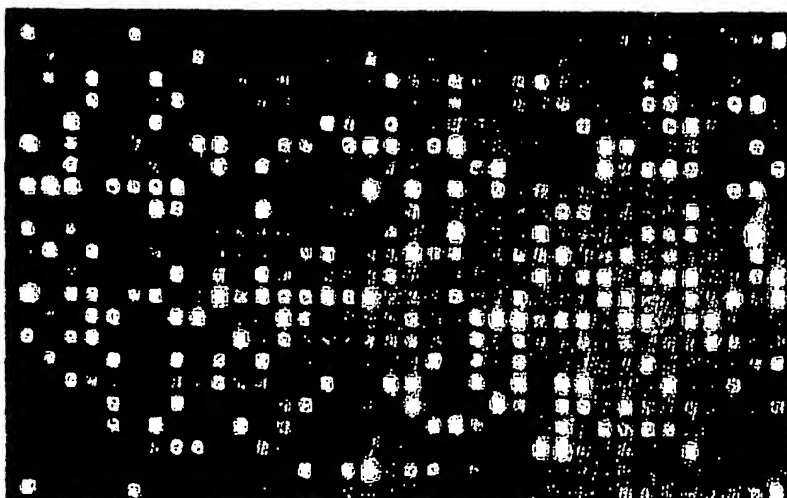


FIG. 4C